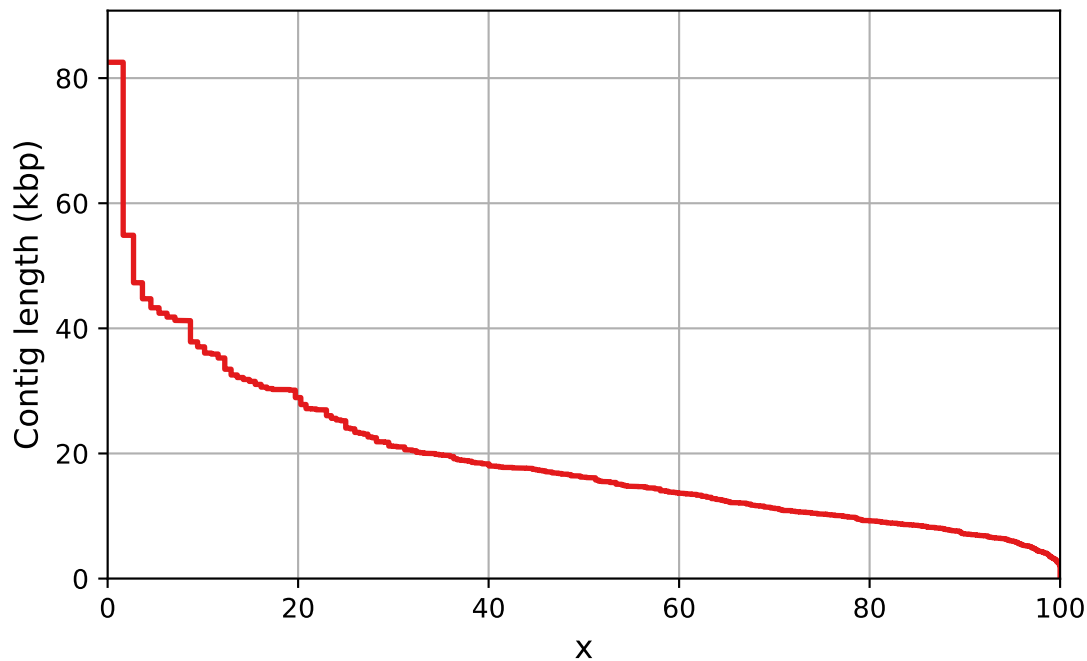


Report

	genome.nextpolish
# contigs (>= 0 bp)	384
# contigs (>= 1000 bp)	384
# contigs (>= 5000 bp)	343
# contigs (>= 10000 bp)	211
# contigs (>= 25000 bp)	36
# contigs (>= 50000 bp)	2
Total length (>= 0 bp)	5052492
Total length (>= 1000 bp)	5052492
Total length (>= 5000 bp)	4901447
Total length (>= 10000 bp)	3896836
Total length (>= 25000 bp)	1263057
Total length (>= 50000 bp)	137409
# contigs	384
Largest contig	82532
Total length	5052492
GC (%)	40.14
N50	16199
N90	7140
auN	19565.4
L50	103
L90	287
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 1 part

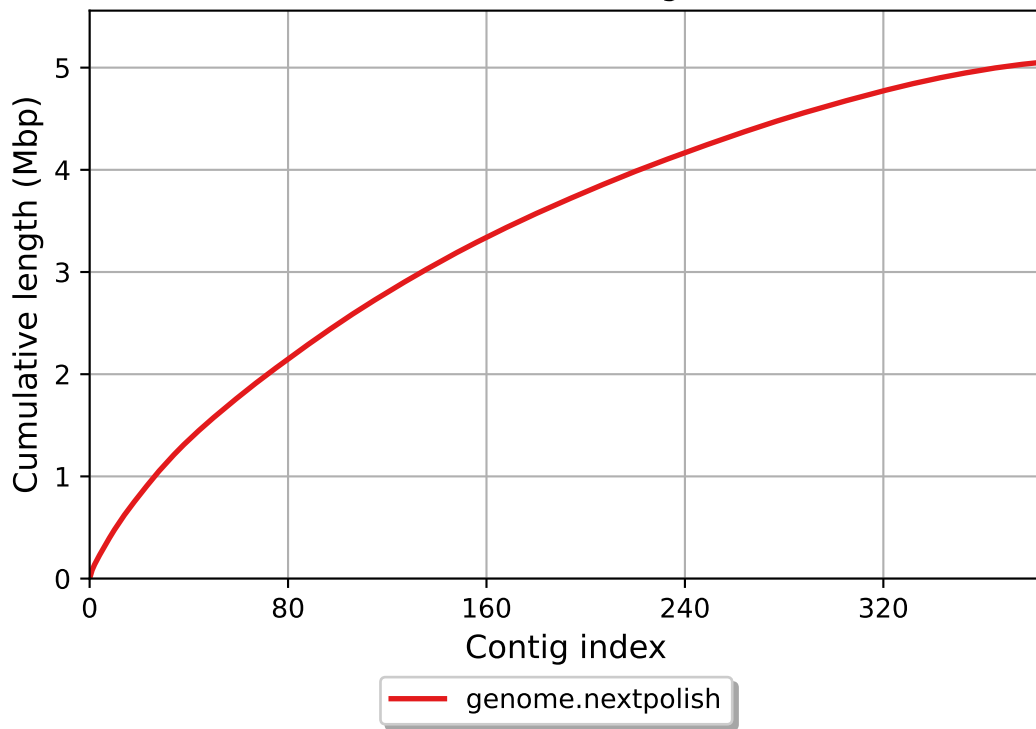
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

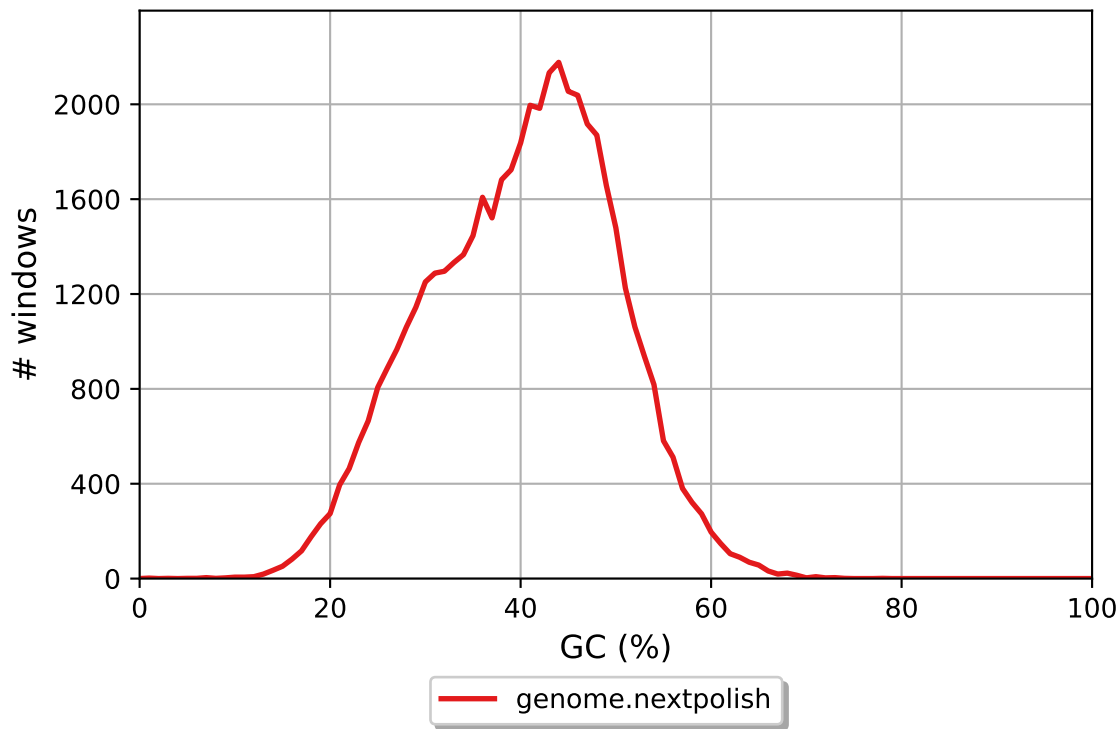


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

